COPY

SEQUENCE LISTING

1

<110> Itoh, Nobuyuki Kavanaugh, W. Michael

<120> HUMAN FGF-21 GENE AND GENE EXPRESSION PRODUCTS

<130> PP-16758.001/201130.408

<140> 09/715,805

<141> 2000-11-16

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<211> 659

<212> DNA

<213> Mus musculus

<220>

<221> CDS

<222> (14)...(646)

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1 5 10

ctg tgg gtc cga ctg ctg ctg gct gtc ttc ctg ctg ggg gtc tac caa 97 Leu Trp Val Arg Leu Leu Leu Ala Val Phe Leu Leu Gly Val Tyr Gln 15 20 25

gca tac ccc atc cct gac tcc agc ccc ctc ctc cag ttt ggg ggt caa 145
Ala Tyr Pro Ile Pro Asp Ser Ser Pro Leu Leu Gln Phe Gly Gln
30 35 40

gtc cgg cag agg tac ctc tac aca gat gac gac caa gac act gaa gcc 193
Val Arg Gln Arg Tyr Leu Tyr Thr Asp Asp Gln Asp Thr Glu Ala
45 50 55 60

cac ctg gag atc agg gag gat gga aca gtg gta ggc gca gca cac cgc 241
His Leu Glu Ile Arg Glu Asp Gly Thr Val Val Gly Ala Ala His Arg
70 75

agt cca gaa agt ctc ctg gag ctc aaa gcc ttg aag cca ggg gtc att 289 Ser Pro Glu Ser Leu Leu Glu Leu Lys Ala Leu Lys Pro Gly Val Ile 80 85 90

caa atc ctg ggt gtc aaa gcc tct agg ttt ctt tgc caa cag cca gat 337 Gln Ile Leu Gly Val Lys Ala Ser Arg Phe Leu Cys Gln Gln Pro Asp

	130					135					140					
Leu 145	Arg	Leu	Pro	Gln	Lys 150	Asp	Ser	Pro	Asn	Gln 155	Asp	Ala	Thr	Ser	Trp 160	
	Pro	Val	Arg	Phe 165		Pro	Met	Pro	Gly 170		Leu	His	Glu	Pro 175		
Asp	Gln	Ala	Gly 180		Leu	Pro	Pro	Glu 185		Pro	Asp	Val	Gly 190		Ser	
Asp	Pro	Leu 195		Met	Val	Glu	Pro 200		Gln	Gly	Arg	Ser 205		Ser	Tyr	
Ala	Ser 210	130					200					200				
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		gtg Val														98
		gac Asp														146
		ctc Leu														194
		gag Glu 65														242
		ctg Leu														290
gga Gly 95	gtc Val	aag Lys	aca Thr	tcc Ser	agg Arg 100	ttc Phe	ctg Leu	tgc Cys	cag Gln	cgg Arg 105	cca Pro	gat Asp	ggg Gly	gcc Ala	ctg Leu 110	338
tat Tyr	gga Gly	tcg Ser	ctc Leu	cac His 115	ttt Phe	gac Asp	cct Pro	gag Glu	gcc Ala 120	tgc Cys	agc Ser	ttc Phe	cgg Arg	gag Glu 125	ctg Leu	386

1)

															ggc Gly		434
															gca Ala		482
															gca Ala		530
	_									_			_	-	ggc Gly		578
															ccc Pro 205		626
		gct Ala		tga *	agco	ca											643
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	Met 1	Asp	Ser	Asp	Glu 5	Thr	Gly	Phe	Glu	His 10	Ser	Gly	Leu	Trp	Val 15	Ser	
				20					25	_				30	Ile		
	Asp	Ser	Ser 35	Pro	Leu	Leu	Gln	Phe 40	Gly	Gly	Gln	Val	Arg 45	Gln	Arg	Tyr	
	Leu	Tyr 50	Thr	Asp	Asp	Ala	Gln 55		Thr	Glu	Ala	His 60	Leu	Glu	Ile	Arg	
	Glu 65		Gly	Thr	Val	Gly 70		Ala	Ala	Asp	Gln 75		Pro	Glu	Ser		
		Gln	Leu	Lys		-	Lys	Pro	Gly			Gln	Ile	Leu	Gly	80 Val	
	Lys	Thr	Ser		85 Phe	Leu	Cys	Gln		90 Pro	Asp	Gly	Ala		95 Tyr	Gly	
	Ser	Leu	His	100 Phe	Asp	Pro	Glu	Ala	105 Cys	Ser	Phe	Arg	Glu	110 Leu	Leu	Leu	
	Glu	Asp	115 Gly	Tyr	Asn	Val	Tyr	120 Gln	Ser	Glu	Ala	His	125 Gly	Leu	Pro	Leu	
		130					135					140			Arg		
	145					150					155				_	160	
	Pro	Ala	Arg	Phe	Leu 165	Pro	Leu	Pro	Gly	Leu 170	Pro	Pro	Ala	Leu	Pro 175	Glu	

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The state of the s
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<213> Mus musculus

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Pro Pro Gly Ile Leu Ala Pro Gln Pro Pro Asp Val Gly Ser Ser Asp
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Pro Leu Ser Met Val Gly Pro Ser Gln Gly Arg Ser Pro Ser Tyr Ala
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Ser
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<213> Homo sapiens

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Gln Ser Val Ser Asp Glu Asp Pro Leu Phe Leu Tyr Gly Trp Gly Lys
Ile Thr Arg Leu Gln Tyr Leu Tyr Ser Ala Gly Pro Tyr Val Ser Asn
Cys Phe Leu Arg Ile Arg Ser Asp Gly Ser Val Asp Cys Glu Glu Asp
                    70
                                        75
Gln Asn Glu Arg Asn Leu Leu Glu Phe Arg Ala Val Ala Leu Lys Thr
                85
                                    90
Ile Ala Ile Lys Asp Val Ser Ser Val Arg Tyr Leu Cys Met Ser Ala
                                105
Asp Gly Lys Ile Tyr Gly Leu Ile Arg Tyr Ser Glu Glu Asp Cys Thr
                            120
Phe Arg Glu Glu Met Asp Cys Leu Gly Tyr Asn Gln Tyr Arg Ser Met
                        135
                                            140
Lys His His Leu His Ile Ile Phe Ile Gln Ala Lys Pro Arg Glu Gln
                    150
                                        155
Leu Gln Asp Gln Lys Pro Ser Asn Phe Ile Pro Val Phe His Arg Ser
                165
                                    170
Phe Phe Glu Thr Gly Asp Gln Leu Arg Ser Lys Met Phe Ser Leu Pro
                                185
Leu Glu Ser Asp Ser Met Asp Pro Phe Arg Met Val Glu Asp Val Asp
                            200
His Leu Val Lys Ser Pro Ser Phe Gln Lys
    210
                        215
<210> 10
<211> 216
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```
Asp Gly Tyr Asn Val Tyr Arg Ser Glu Lys His Arg Leu Pro Val Ser
                              135
                                                   140
     Leu Ser Ser Ala Lys Gln Arg Gln Leu Tyr Lys Asn Arg Gly Phe Leu
                          150
                                               155
     Pro Leu Ser His Phe Leu Pro Met Leu Pro Met Val Pro Glu Glu Pro
                                           170
     Glu Asp Leu Arg Gly His Leu Glu Ser Asp Met Phe Ser Ser Pro Leu
                                       185
     Glu Thr Asp Ser Met Asp Pro Phe Gly Leu Val Thr Gly Leu Glu Ala
             195
                                  200
     Val Arg Ser Pro Ser Phe Glu Lys
         210
     <210> 11
     <211> 10
     <212> PRT
     <213> Unknown
l, k
     <220>
L
     <223> Residues which contain the anitgenic determinant
THE WAY
           recognized by the myc monoclonal antibody.
Marie House
     <400> 11
     Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
ij1
W.
E.
<210> 12
     <211> 5
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The first
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     <223> Preferred thrombin cleave site.
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                       5
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      <212> PRT
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      <400> 14
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     <211> 6
     <212> PRT
     <213> Homo sapiens
     <400> 15
     Leu Pro Met Leu Pro Met
      1
                         5
     <210> 16
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     <213> Homo sapiens
THE STATE OF
E.
     <400> 16
    Arg Pro Asp Gly Tyr Asn 1 5
#
The start that they done in the
    <210> 17
    <211> 6
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    <213> Homo sapiens
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    His Phe Leu Pro Met Leu
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